

CRF Errors Corrected by the STIC Systems Branch.

Serial Number: 09/913,351A

CRF Processing Date:

Edited by:

Verified by:

5/20/2003

(STIC staff)

ENTERED Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped" down to the next line. Edited a format error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____ Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as _____ Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____ Other:

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1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/913,351A

DATE: 05/20/2003

TIME: 08:36:51

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\05192003\I913351A.raw

3 <110> APPLICANT: DELGADO, AURORA BRIEVA
 4 VILLARRUBIA, VINCENTE GARCIA
 5 GOMEZ-PAMO, ANTONIO GUERRERO
 6 • RANIERI, JUAN PABLO PIVEL
 7 GALLEGOS, GUILLERMO GIMENEZ
 8 TUDURI, JOSE ANTONIO MATJI
 10 <120> TITLE OF INVENTION: PHARMACOLOGICALLY ACTIVE POLYPEPTIDE GLYCOCONJUGATES
 12 <130> FILE REFERENCE: 618999-1/JP/B-4275
 14 <140> CURRENT APPLICATION NUMBER: 09/913,351A
 15 <141> CURRENT FILING DATE: 1999-10-21
 17 <150> PRIOR APPLICATION NUMBER: PCT/ES99/00338
 18 <151> PRIOR FILING DATE: 1999-10-21
 20 <150> PRIOR APPLICATION NUMBER: ES P9900408
 21 <151> PRIOR FILING DATE: 1999-02-26
 23 <160> NUMBER OF SEQ ID NOS: 5
 25 <170> SOFTWARE: PatentIn Ver. 2.1
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 230
 29 <212> TYPE: PRT
 30 <213> ORGANISM: Artificial Sequence
 32 <220> FEATURE:
 33 <223> OTHER INFORMATION: Description of Artificial Sequence: Formula
 34 sequence
 36 <220> FEATURE:
 37 <221> NAME/KEY: MOD_RES
 38 <222> LOCATION: (1)..(48) ✓
 39 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
 40 3-48 amino acids
 42 <220> FEATURE:
 43 <221> NAME/KEY: MOD_RES
 44 <222> LOCATION: (50)..(62)
 45 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
 46 9-13 amino acids
 48 <220> FEATURE:
 49 <221> NAME/KEY: MOD_RES
 50 <222> LOCATION: (64) ✓
 51 <223> OTHER INFORMATION: Gln, Glu, Arg or Lys
 53 <220> FEATURE:
 54 <221> NAME/KEY: MOD_RES
 55 <222> LOCATION: (65) ✓
 56 <223> OTHER INFORMATION: Variable amino acid
 59 <220> FEATURE:
 60 <221> NAME/KEY: MOD_RES

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62 <223> OTHER INFORMATION: Hydrophobic amino acid
64 <220> FEATURE:
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66 <222> LOCATION: (67)
67 <223> OTHER INFORMATION: Leu, Ile, Val or Met
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70 <221> NAME/KEY: MOD_RES
71 <222> LOCATION: (68)..(106)
72 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
73 15-39 amino acids
75 <220> FEATURE:
76 <221> NAME/KEY: MOD_RES
77 <222> LOCATION: (109)
78 <223> OTHER INFORMATION: Hydrophilic amino acid
80 <220> FEATURE:
81 <221> NAME/KEY: MOD_RES
82 <222> LOCATION: (110)
83 <223> OTHER INFORMATION: Gln, Glu or His
85 <220> FEATURE:
86 <221> NAME/KEY: MOD_RES
87 <222> LOCATION: (111)
88 <223> OTHER INFORMATION: Leu or Val
90 <220> FEATURE:
91 <221> NAME/KEY: MOD_RES
92 <222> LOCATION: (112)..(117)
93 <223> OTHER INFORMATION: Variable amino acid
95 <220> FEATURE:
96 <221> NAME/KEY: MOD_RES
97 <222> LOCATION: (119)
98 <223> OTHER INFORMATION: Variable amino acid
100 <220> FEATURE:
101 <221> NAME/KEY: MOD_RES
102 <222> LOCATION: (121)..(122)
103 <223> OTHER INFORMATION: Variable amino acid
105 <220> FEATURE:
106 <221> NAME/KEY: MOD_RES
107 <222> LOCATION: (123)
108 <223> OTHER INFORMATION: Leu or Ile
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112 <222> LOCATION: (124)..(179)
113 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
114 13-56 amino acids
117 <220> FEATURE:
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119 <222> LOCATION: (181)..(206)
120 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
121 15-26 amino acids

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124 <221> NAME/KEY: MOD_RES
125 <222> LOCATION: (208)
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128 <220> FEATURE:
129 <221> NAME/KEY: MOD_RES
130 <222> LOCATION: (209)
131 <223> OTHER INFORMATION: Val, Ile, Leu or Met
133 <220> FEATURE:
134 <221> NAME/KEY: MOD_RES
135 <222> LOCATION: (210)...(217)
136 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
137 1-8 amino acids
139 <220> FEATURE:
140 <221> NAME/KEY: MOD_RES
141 <222> LOCATION: (218)...(230)
142 <223> OTHER INFORMATION: Variable amino acid; this Xaa range may encompass
143 1-12 amino acids
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147 1 5 10 15
149 Xaa Xaa
150 20 25 30
152 Xaa Xaa
153 35 40 45
155 Cys Xaa Cys Xaa
156 50 55 60
158 Xaa Xaa
159 65 70 75 80
161 Xaa Xaa
162 85 90 95
164 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Xaa Xaa Xaa Xaa
165 100 105 110
167 Xaa Xaa Xaa Xaa Cys Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
168 115 120 125
170 Xaa Xaa
171 130 135 140
173 Xaa Xaa
174 145 150 155 160
176 Xaa Xaa
177 165 170 175
179 Xaa Xaa Xaa Gly Xaa Xaa
180 180 185 190
182 Xaa Cys Xaa
183 195 200 205
185 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
186 210 215 220
188 Xaa Xaa Xaa Xaa Xaa Xaa
189 225 230

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RAW SEQUENCE LISTING

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TIME: 08:36:51

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\05192003\I913351A.raw

192 <210> SEQ ID NO: 2
193 <211> LENGTH: 37
194 <212> TYPE: PRT
195 <213> ORGANISM: Ricinus communis
197 <400> SEQUENCE: 2
198 Glu Ser Lys Gly Glu Arg Glu Gly Ser Ser Gln Gln Cys Arg Gln
199 1 5 10 15
201 Glu Val Gln Arg Lys Asp Leu Ser Ser Cys Glu Arg Tyr Leu Arg Gln
202 20 25 30
204 Ser Ser Ser Arg Arg
205 35
208 <210> SEQ ID NO: 3
209 <211> LENGTH: 68
210 <212> TYPE: PRT
211 <213> ORGANISM: Ricinus communis
213 <400> SEQUENCE: 3
214 Gln Gln Gln Glu Ser Gln Gln Leu Gln Gln Cys Cys Asn Gln Val Lys
215 1 5 10 15
217 Gln Val Arg Asp Glu Cys Gln Cys Glu Ala Ile Lys Tyr Ile Ala Glu
218 20 25 30
220 Asp Gln Ile Gln Gln Gly Gln Leu His Gly Glu Glu Ser Glu Arg Val
221 35 40 45
223 Ala Gln Arg Ala Gly Glu Ile Val Ser Ser Cys Gly Val Arg Cys Met
224 50 55 60
226 Arg Gln Thr Arg
227 65
230 <210> SEQ ID NO: 4
231 <211> LENGTH: 34
232 <212> TYPE: PRT
233 <213> ORGANISM: Ricinus communis
235 <400> SEQUENCE: 4
236 Pro Ser Gln Gln Gly Cys Arg Gly Gln Ile Gln Glu Gln Gln Asn Leu
237 1 5 10 15
239 Arg Gln Cys Gln Glu Tyr Ile Lys Gln Gln Val Ser Gly Gln Gly Pro
240 20 25 30
242 Arg Arg
246 <210> SEQ ID NO: 5
247 <211> LENGTH: 65
248 <212> TYPE: PRT
249 <213> ORGANISM: Ricinus communis
251 <400> SEQUENCE: 5
252 Gln Glu Arg Ser Leu Arg Gly Cys Cys Asp His Leu Lys Gln Met Gln
253 1 5 10 15
255 Ser Gln Cys Arg Cys Glu Gly Leu Arg Gln Ala Ile Glu Gln Gln Gln
256 20 25 30
258 Ser Gln Gly Gln Leu Gln Gly Gln Asp Val Phe Glu Ala Phe Arg Thr
259 35 40 45
261 Ala Ala Asn Leu Pro Ser Met Cys Gly Val Ser Pro Thr Glu Cys Arg
262 50 55 60

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TIME: 08:36:51

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\05192003\I913351A.raw

264 Phe
265 65

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/913,351A

DATE: 05/20/2003
TIME: 08:36:52

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\05192003\I913351A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22.
Seq#:1; Xaa Pos. 23,24,25,26,27,28,29,30,31,32,33,34,35,36,37,38,39,40,41
Seq#:1; Xaa Pos. 42,43,44,45,46,47,48,50,51,52,53,54,55,56,57,58,59,60,61
Seq#:1; Xaa Pos. 62,64,65,66,67,68,69,70,71,72,73,74,75,76,77,78,79,80,81
Seq#:1; Xaa Pos. 82,83,84,85,86,87,88,89,90,91,92,93,94,95,96,97,98,99,100
Seq#:1; Xaa Pos. 101,102,103,104,105,106,109,110,111,112,113,114,115,116
Seq#:1; Xaa Pos. 117,119,121,122,123,124,125,126,127,128,129,130,131,132
Seq#:1; Xaa Pos. 133,134,135,136,137,138,139,140,141,142,143,144,145,146
Seq#:1; Xaa Pos. 147,148,149,150,151,152,153,154,155,156,157,158,159,160
Seq#:1; Xaa Pos. 161,162,163,164,165,166,167,168,169,170,171,172,173,174
Seq#:1; Xaa Pos. 175,176,177,178,179,181,182,183,184,185,186,187,188,189
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Seq#:1; Xaa Pos. 204,205,206,208,209,210,211,212,213,214,215,216,217,218
Seq#:1; Xaa Pos. 220,221,222,223,224,225,226,227,228,229,230